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			10		20		30		40
			TPVSEKQLAE	VVANTITPLM	KAQSVPGMAV	AVIYQ GKPHY			
50	60	70		80	90	100			
YTFGKADIAA	NKPVTPQTLF	ELGSISKTF	* *	GVLGGDAIAR	GEISLDDAVT	RYWPQLTGKQ			
110	120	130		140	150	160			
WQGIRMLDLA	TYTAGGLPLQ	VPDEVTDNAS		LLRFYQNWQP	QWKPGTTRLY	ANASIGLFGA	*	*	
170	180	190		200	210	220			
LAVKPSGMPY	EQAMTTRVLK	PLKLDHTWIN		VPKAEAAHYA	WGYRDGKAVR	VSPGMLDAQA			
230	240	250		260	270	280			
YGVKTNVQDM	ANWVMANMAP	ENVADASLKQ		GIALAQSRYW	RIGSMYQGLG	WEMLNWPVEA			
290	300	310		320	330	340			
NTVVEGSDSK	VALAPLPVAE	VNPPAPPVKA	** *	SWVHKTGSTG	GFGSYVAFIP	EKQIGIVMLA			
350	360								
NTSYPNPARV	EAAYHILEAL	Q							

FIGURE 1

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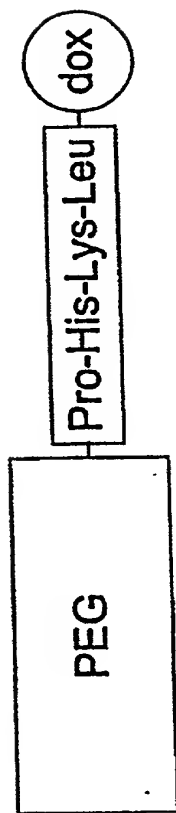


FIGURE 2

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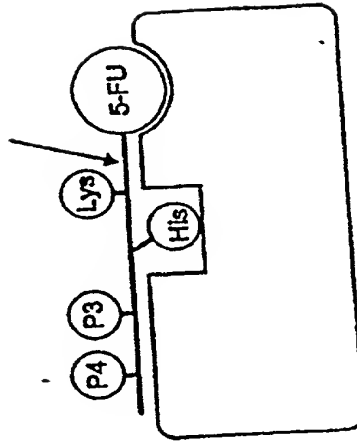


FIGURE 3

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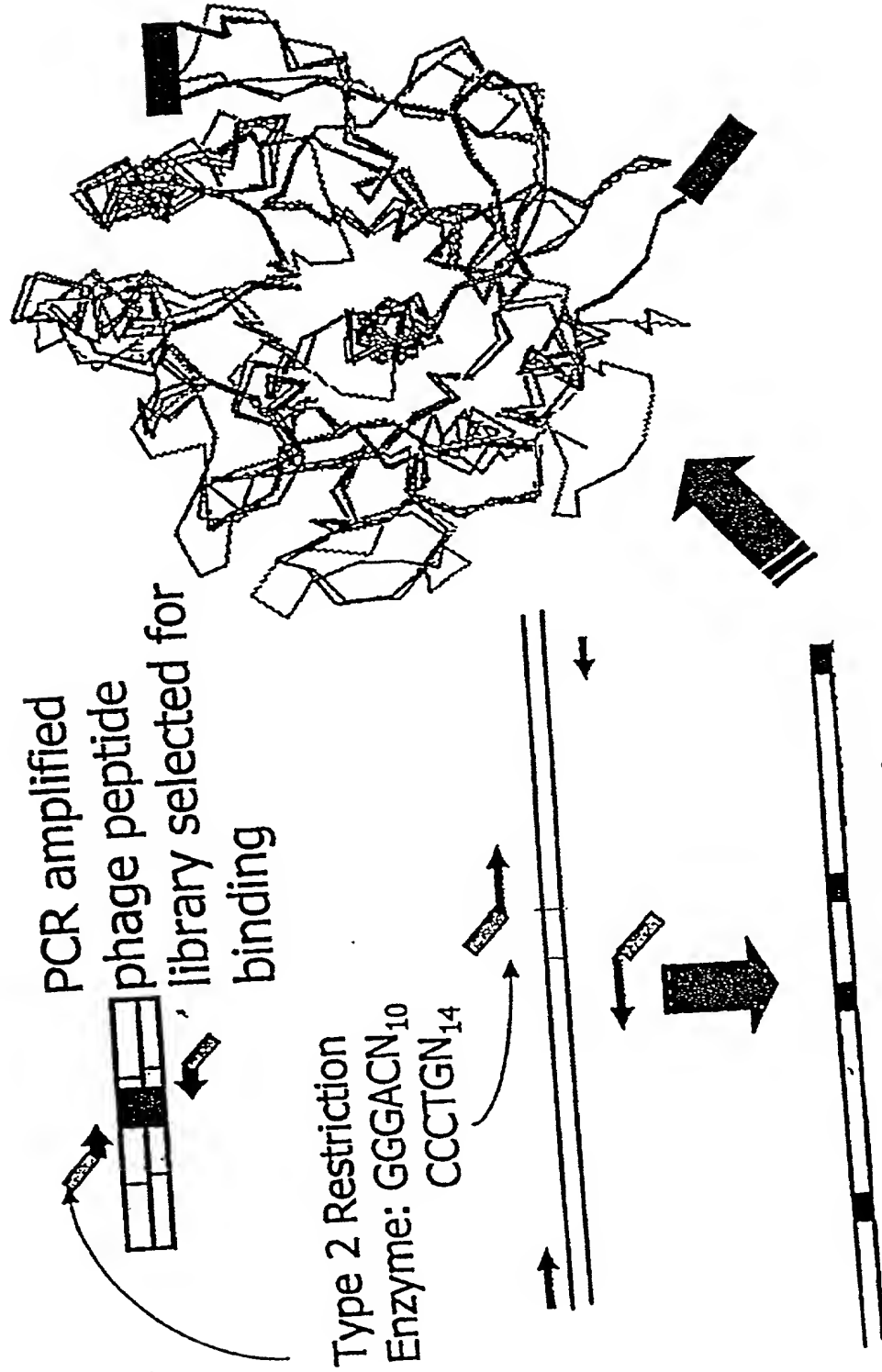


FIGURE 4

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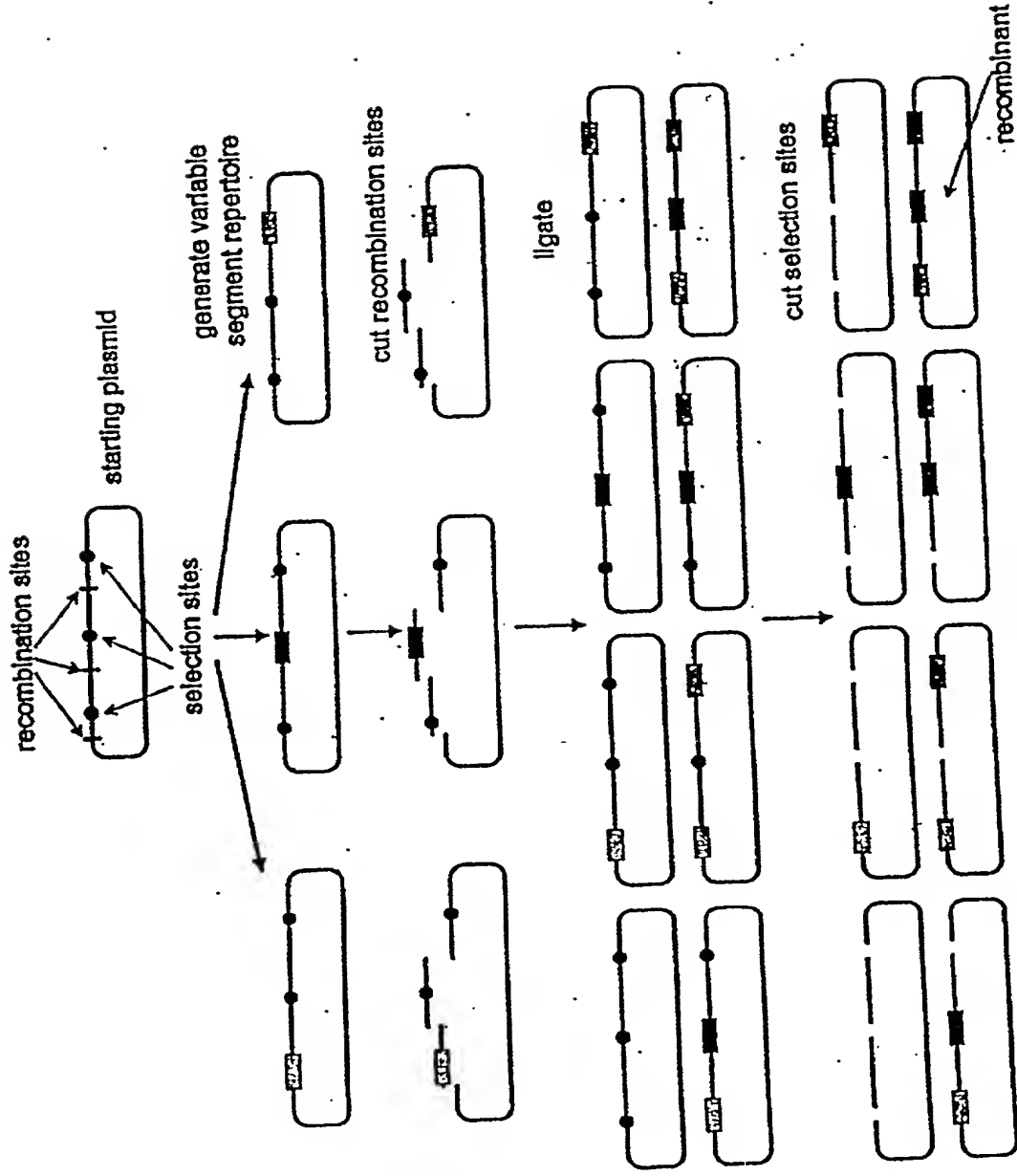


FIGURE 5

FIGURE 6

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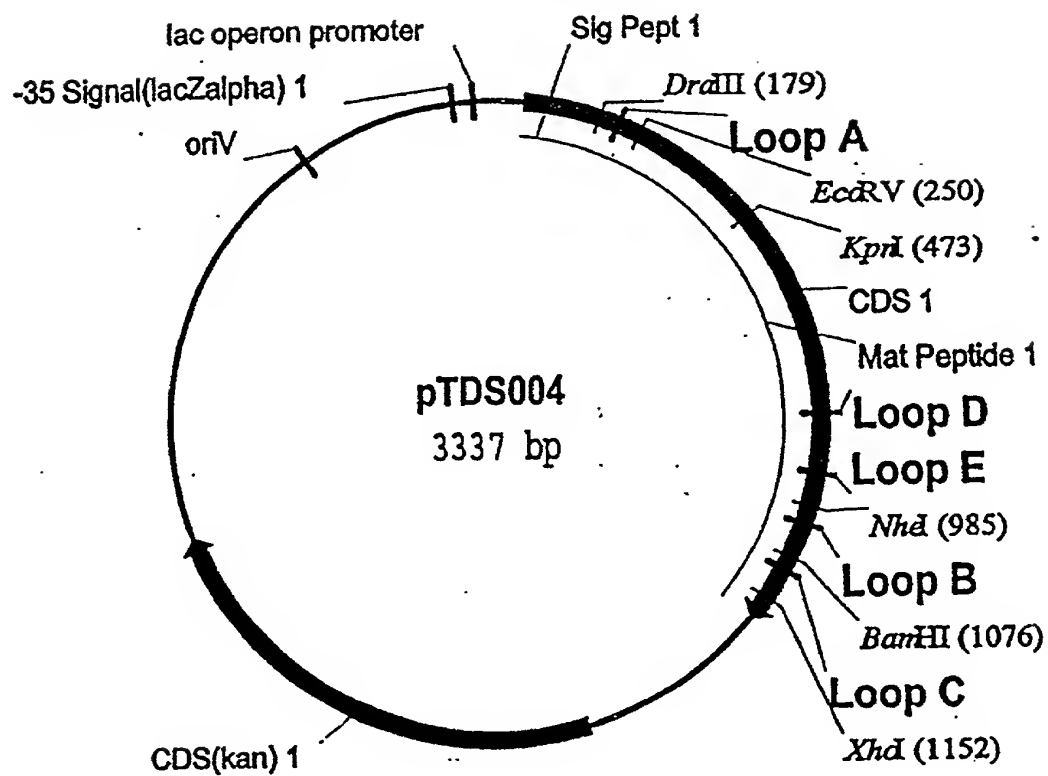


FIGURE 7

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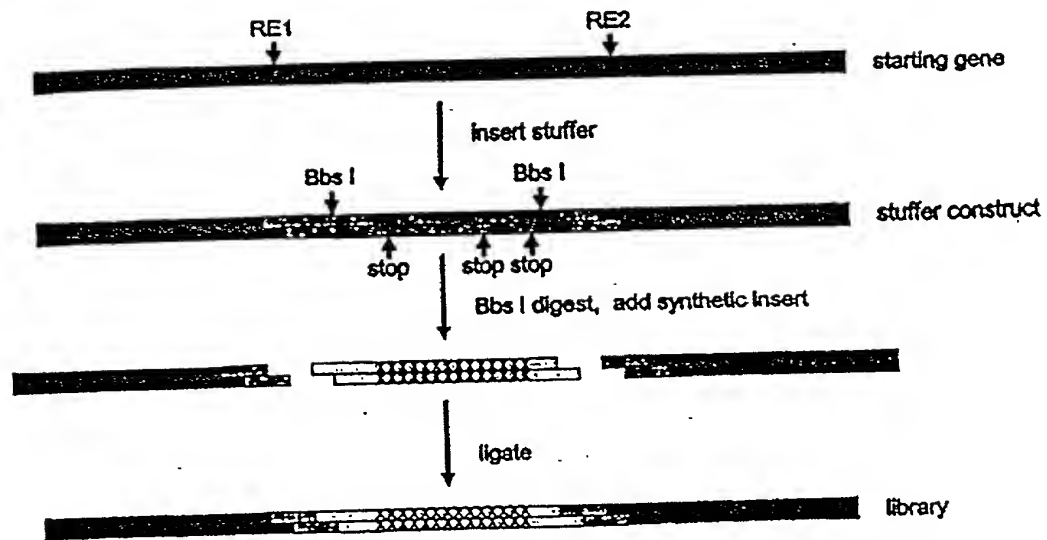


FIGURE 8

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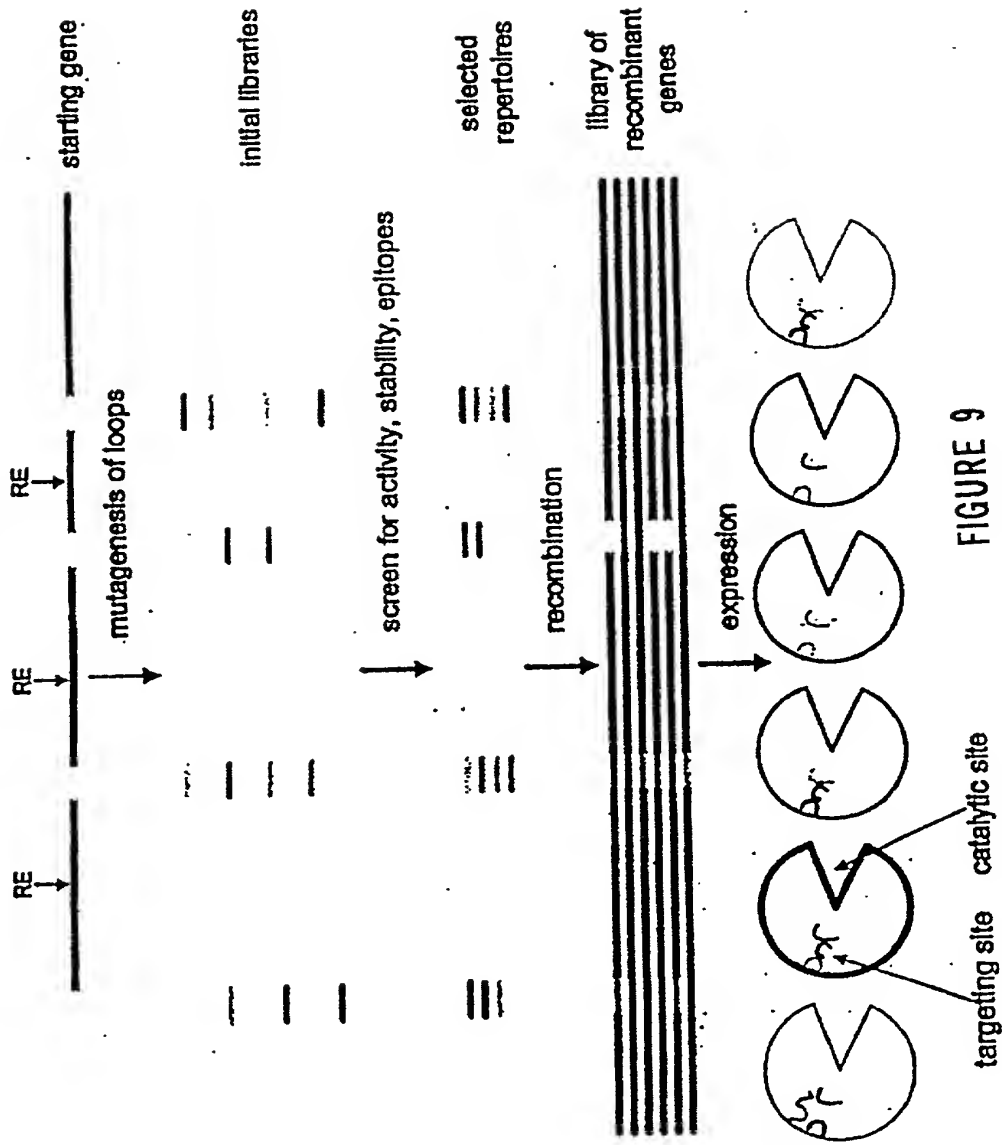


FIGURE 9

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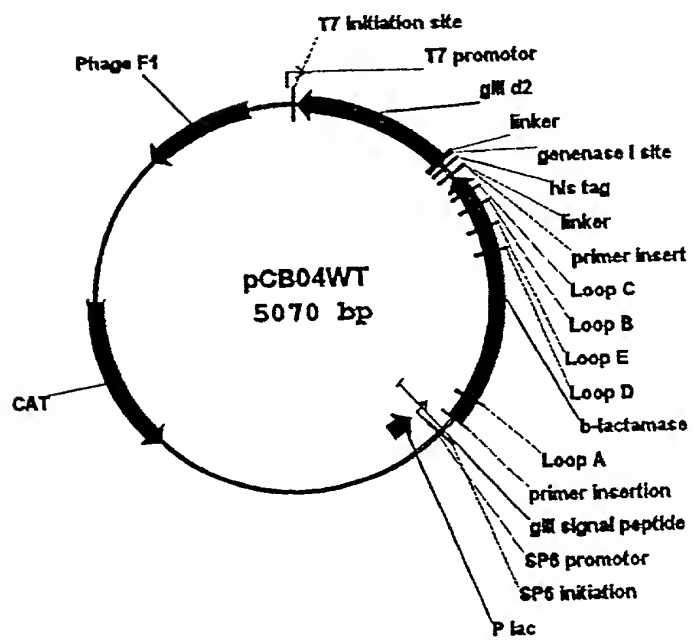


FIGURE 10